Bibliographic

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Title:
Perfect score:
Sequence:
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Listing first
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RESULT 1
AR133394 15 bp DNA linear PAT 16-MAY-2001
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LOCUS 394
DEFINITION Sequence 1819 from patent US 6194150.
ARCESSION AR133394
VERSION AR133394
VERSION AR133394.1 GI:14122299
KEYWORDS 5URCE Unknown.
ORGANISM Unknown.
Unclassified.
TITLE 1 (bases 1 to 15)
AUTHORS 5tinchcomb,D.T., Jarvis,T. and McSwiggen,J.
FEATURES 5tinchcomb,D.T.
FEATURES 5tinchcom
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Query Match Best Local Similarity

100.0%;

Score Pred.

No.

DB 6; L, 1.2e+05;

Length 15;

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RESULT 4
AR043730/c
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The high bone mass gene of 11q13.3
Patent: WO 0177327-A 32 18-OCT-2001;
Genome Therapeutics Corporation (US)
Location/Qualifiers
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AX277553
AX277553.1 GI:16604752
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Nucleic acid based inhibition of CD40
Patent: US 6194150-A 1820 27-FEB-2001;
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Sequence 1820 from patent
AR133395
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/note="Artificial sequence is a primer."
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Sequence 101 from patent US
AR043731
AR043731.1 GI:5964739
1 (bases 1 to 19) Robbins, J.M. and T. Ribozyme therapy for
                                                       Homo sapiens
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                                 Eukaryota; Metazoa; Chordata;
Mammalia; Eutheria; Primates;
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Seidel, H. Martin., Lamb, I. Peter. and Chan, S.-S. Tian. Methods for detecting modulators of cytokine action Patent: US 5712094-A 56 27-JAN-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 56 from I81958 I81958.1 GI:3210
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Patent: WO 0130362-A 1530 03-MAY-2001;
IMMUSOL, INC. (US)
                                                                                 Unclassified.

1 (bases 1 to 19)
Seidel, H. Martin., Lamb, I. Peter. and Chan, S. - S. Tian. Methods for detecting modulators of cytokine action Patent: US 5712094-A 57 27-JAN-1998;
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/db_xref="taxon:9606"
/note="Cyclin A2 ribozyme
2 c 5 g 4
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Sequence 21
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La Thangue, N.B. and de la Luna, S.
Transcription factor DP-3 and isoforms
Patent: US 5859199-A 21 12-JAN-1999;
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La,T.N. and De,L.L.
DP AND E2F PROTEIN NUCLEAR LOCALISATION
Patent: WO 9743647-A 21 20-NOV-1997;
MEDICAL RES COUNCIL (GB)
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Sequence 21 from Patent W09743647.
A67540 GI:4756420
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LE PATENT: JP 1992058888-A 2 25-FEB-1992;

SHIONOGI & CO LTD

OS Artificial sequence; Genes.
PN JP 1992058888-A/2
PD 25-FEB-1992
PC C12N15/11,C07H21/00,C12Q1/68,G01N33/569,G01N33/574; CC strandedness: Single; CC topology: Linear; CC topology: Linear; CC anti-sense: No; CC anti-sense: No;
                                              TTCTGAGAA 12
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Oligonucleotide for detection
E03365 GI:2171582
JP 1992058888-A/2.
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La Thangue, N.D. and De La Luna, S.
Assay for a putative regulator of cell
Patent: US 6159691-A 21 12-DEC-2000;
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1 (bases 1 to 22)

Stemmer, W.P.C.

Methods of shuffling patent: US 6132970-A:
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Sequence 18
AR051826
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Stemmer,W.P.C. and Crameri,A.
DNA mutagenesis by random fragmentation
Patent: US 5830721-A 18 03-NOV-1998;
Location/Qualifiers
1...22
                                                                                       Sequence 18 from patent
AR115962
AR115962.1 GI:14096284
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Stemmer,W.P.C. and Crameri,A.
Methods for generating polynucleotides having desired characteristics by iterative selection and recombination patent: US 5811238-A 18 22-SEP-1998;
Location/Qualifiers
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Sequence 18 from patent
AR040919
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ALIGNMENTS

AAT14149 standard; DNA; 9

ВP

AAT14149;

Cytokine responsive DNA spacer regulatory element.

29-MAY-1996 (first entry)

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RESULT 1
AAT14149/c
ID AAT141
XX AAT141
XX AAT141
XX AAT141
XX PEGULA
XX REGULA
XX REGULA
XX SIGNAL
XX Gene t
XX Gene t
XX Gene
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XX 10-APR
XX 11-APR
XX 17-MAR
PR 14-APR
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XX 11-APR

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DNA spacer regulatory elements responsive to cytokine(s) -
                                                                                               WPI; 1995-373797/48.
                                                                                                                                                                                                                                                                                          (LIGA-) LIGAND PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                         27-MAR-1995;
14-APR-1994;
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94US-0228935.
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RESULT 2
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XX Regula
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XX Synthe
YN W09630
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Matches 9
                       a regulatory element having the present sequence, operably linked to a promoter, which is operably linked to a heterologous gene (preferably a marker gene). The gene is under the transcriptional control of the promoter and the ON sequence when the ON is bound a STAT protein activated in response to II-3, II-3, G-CSF, GM-CSF erythropoietin, thrombopoletin, or preferably II-4, II-7, II-9, II-13 or II-15. Cytokine responsive host cells transfected with t
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           spacer regulatory exemient on a sample, and in assays a transcriptional regulatory protein in a sample, and in assays for (ant)agonists of gene transcription. The identified cpds. may be used to treat cytokine-induced disease states, or to ameliorate disease states caused by cytokine deficiency, e.g. and in figure and in the conditions of the conditions and in the conditions of the conditions are conditions.
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                                                                                                                                                                                                        A novel DNA construct comprises an oligonucleotide (ON)
                                                                                                                                                                                                                                                                                                         DNA construct for screening modulators of cytokine-mediated transcription - contg. regulatory element and a cytokine-sensitive promoter operably linked to a heterologous gene
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                                                                                                                                                                                                                                                               Claim 5;
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Matches
                                                              AAV56842-V56976 and AAV61601-V61631 are oligonucleotides used in the production of constructs comprising a cytckine-responsive regulatory element linked to a promoter which is linked to a heterologous coding sequence so that the coding sequence is under the transcriptional control of the regulatory element and the promoter, where the regulatory element has a nucleotide sequence selected from TTONNGAA, TTANYTAA, and TTCNYTAA where N is A, T, C or G, and y = 3 or 4. The constructs can be used to detect or isolate transcription-regulating proteins, e.g. STAT proteins, in a sample by contacting the sample with the construct so that the protein binds to the regulatory element, and detecting or separating the resulting complex. The cells can be used in screening assays for
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coding se
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             act as an agonist or antagonist of cytokine mediated gene transcription. In particular, they can be used to screen cytokine modulators involved in the STAT5 and/or STAT6 presignalling pathway.
                                                                                                                                                                                                                                                                           DNA constructs containing cytokine-responsive regulatory elements useful in assays for transcription-regulating proteins or gene
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ilarity 100.0%;
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94US-0228935.
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                        transcription, in which the level of expression of the is measured in the presence and absence of a test he presence of the corresponding cytokine.
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The present invention describes a novel enzymatic nucleic acid (ENA) CC having a hammerhead motif (HM) comprising: (1) at least 5 ribose cresidues; (ii) a 2'-C-allyl modification at position 4 of the ENA; (iii) at least ten 2'-O-methyl modifications; and (iv) a 3'-end modification. CC The ENA's can inhibit collagenase and stromelysin production in the synovial membrane of joints for the treatment or prevention of arthritis. CC particularly osteoarthritis or rheumatoid arthritis. The ENA's can also be used to treat antigen presenting cells of a donor to induce tolerance in a recipient to an alloantigen of a donor. They can also be used for cenhancing graft tolerance or for treating alloridations. The ENA's can also be used in diagnosis. Ribozyme therapy impacts on the expression of expression without introducing the non-specific effects upon gene expression which accompany treatment with retinoids and dexamethasone. The concentration of ribozyme required to affect a therapeutic treatment cc is lower than that required of antisense molecules, and is highly
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                                                                                                                                                                                                               Claim
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for the to
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                                                                                                                                                                                                                                                  ymatic nucleic acid molecules having a the treatment of arthritis, induction
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                                                                                                                                                                                                                                                                                                                          Gustofson J,
L, Karpeisky
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A, Modak A,
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Wincott
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cott F;
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                                                                                                                                                                                                        Enzymatic nucleic acid molecules having for the treatment of arthritis, inductive treatment of auto-immune diseases
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Arthritic condition; graft tolerance; immune response; target; cl hammerhead ribozyme; hairpin ribozyme; human; rabbit; mouse; coll stromelysin; synovial membrane; joint; arthritis; osteoarthritis;
                                                                                                                                                                                                                                                                                                                                           Draper K, G
Beigelman L,
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                                                                                                                                                                                                                                                                                                                                                             Gustofson J,
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c J, Jarvis
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94US-0363254.
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A, Modak A, Usman
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The present invention describes a novel enzymatic nucleic acid (ENA) having a hammerhead motif (HM) comprising: (1) at least 5 ribose residues; (11) a 2'-C-allyl modification at position 4 of the ENA; (111) at least ten 2'-O-methyl modifications; and (1v) a 3'-end modification. The ENA's can inhibit collagenase and stromelysin production in the

Claim

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Page 198;

307pp;

English.

The ENA's can inhibit collagenase and stromelysin production in the synovial membrane of joints for the treatment or prevention of arthritis, particularly osteoarthritis or rheumatoid arthritis. The ENA's can also be used to treat antigen presenting cells of a donor to induce tolerance

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                                                                                 The present invention relates to enzymatic and antisense nucleic molecules that act as inhibitors of the expression of repressor geneding the TR2 Orphan receptor. EAR3/COUP-TF-1, the GARTA transcription factor gene, IRF-2 and/or the CAATT Displacement Protein (CDP). Inhibition of the repressors removes prevents inhibition (and consequently increases expression of) genes involute production of erythropoietin, granulocyte colony stimulating protein and interferon alpha.
                                                                                                                                                                                                   Claim
                                                                                                                                                                                                                         Enzymatic and antisense nucleic acid inhibition of repressor genes, useful for producing e.g. granulocyte colony stimulating factor protein, interferon alpha and erythropoletin -
                                                                                                                                                                                                                                                                                                                                                                                                                                                    Homo sapiens.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           present invention
                                                              Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                              interferon alpha;
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                                                                                                                                                                                                   37; Page 76;
  Similarity
9; Conserv
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9; Conser
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tive 0;
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Pred. No. 9.5
0; Mismatches
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AAF02602/c
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                                                                                                                                                                                             Query Match
Best Local 9
                                                                                                                                                                                    Matches
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                                                                              .T 8
1603/c
AAF02603 standard; DNA; 17
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                                                                                                                                                                                                                                                                                                                                                                               protein,
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                      Hammerhead ribozyme substrate #898.
                                                                 AAF02603;
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                                           16-FEB-2001
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| 16 TTCTGAGAA
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                                                                                                                                                                                  Similarity 9; Conserv
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erythropoietin; granulocyte
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                                                                                                                                                                                 100.0%;
ilarity 100.0%;
Conservative (
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3. 9.5e+03;
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Best Local Similarity
Matches 9; Conser
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI;
                                                                                                                                                                                                                                                                                                                                               Human: single nucleotide polymorphism; SNP; genotyping; DNA analysis; allele specific oligonucleotide; ASO; reduced complexity genome; RCG; genomic classification; identification; DNA fingerprinting; tumour characterisation; hybridisation; ss.
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                                                                                                    25-SEP-1998;
                                                                                                                                                   24-SEP-1999;
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                                                                                                                                                                                                                                                                                                 Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human genomic SNP allele specific oligonucleotide
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                                                   (MASI ) MASSACHUSETTS INST TECHNOLOGY.
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     JE,
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  В,
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  Housman DE,
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9.5e+03;
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Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Detection of single nucleotide polymorphisms in genomes by preparation and analysis of reduced complexity genomes, useful for genotyping, fingerprinting and determining allele frequency of SNPs -
                                                                                                                                     Sequence 17
                                                                                                                                                                                                                             to AAA36632
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                                                                                                                                 BP; 4 A; 1 C; 4 G; 8 T; 0 other;
                                                                                                                                                                                                                        represent nucleotide sequences containing SNPs.
    21;
Length 17;
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RESULT 10
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                                                                            (KENO/) IKENO M.
(MASU/) MASUMOTO H.
(OKAZ/) OKAZAKI T.
                                                                                                        26-AUG-1996;
                                                                                                                      26-AUG-1996;
                                                                                                                                    05-MAR-1998
                                                                                                                                                  WO9808964-A1
                                                                                                                                                                        Synthetic.
                                                                                                                                                                                                                 Primer alpha(1)a for human alphoid DNA sequence.
                                                                                                                                                                                                                                12-AUG-1998
                                                                                                                                                                                                                                              AAV24227;
                                                                                                                                                                                                                                                             AAV24227 standard;
                                                                                                                                                                Homo sapiens
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nes 9; Conserv
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Pred. No.
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Mammalian artificial comprising mammalian CENP-B box sequences

telomere and chromosomes -

prepared using DNA construct centromere containing copies of

WPI; 1998-216941/19

Cooke HJ,

Grimes BR,

Ikeno

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Masumoto

Ή,

Okazaki T;

Disclosure; Page

34;

95pp; English

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CC This sequence is a primer for a human alphoid DNA sequence that can be CC used in the DNA construct of the invention. The construct is a mammalian CC artificial chromosome (MAC), and comprises a mammalian telomere and a CC centromere, where the centromere has a DNA sequence containing copies of the CENP-B box sequence. The invention also relates to methods for CC establishing yeast artificial chromosomes (YACs) comprising mammalian CC telomeres and centromeres. The method can be used to construct, modify CC and stably maintain YACs in yeast cells which have the ability to form CC MACs when introduced into mammalian cells. The MACs can replicate CC autonomously, be stably maintained extrachromosomally and transmitted CC efficiently in mammalian cells. The MACs can be used for basic studies on CC organisation and function of mammalian chromosomes and also as a vector to introduce DNA segments (genes) of interest to test their functions in CC mammalian cells or bodies, since the genes carried by MACs will neither CC eause unpredictable insertion mutation on host chromosomes. Furthermore CC where an entire large gene or group of genes and regulatory elements CC could be included. The MACs can be used for somatic gene therapy or to commatte transcapic mice.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Matches
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Best Local
                                                       comprising CENP-B box
                        Disclosure; Page 34; 95pp; English
                                                                                                                                                                                                                                                                                                                         WO9808964-A1
                                                                                                                                                                                                                                                                                                                                                                                                             CENP-B box; mammalian artificial chromosome; MAC; chromosome 21; mammalian telomere; centromere; yeast artificial chromosomes; YAC;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence
                                                                     Mammalian artificial comprising mammalian
                                                                                                                 WPI; 1998-216941/19.
                                                                                                                                              Cooke
                                                                                                                                                                                                                                      26-AUG-1996;
                                                                                                                                                                                                                                                                 26-AUG-1996;
                                                                                                                                                                                                                                                                                               05-MAR-1998
                                                                                                                                                                                                                                                                                                                                                                       Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                               somatic gene therapy; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                          Primer alpha(1)b for human alphoid DNA sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       12-AUG-1998 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAV24228;
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                                                                                                                                                                                         (KENO/) IKENO
(MASU/) MASUMO
                                                                                                                                                                                                                                                                                                                                                      Homo sapiens
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                                                                                                                                              HJ, Grimes BR,
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                                                                                                                                                                        MASUMOTO H.
OKAZAKI T.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      transgenic mice.
                                                       sequences
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                                                                     chromosomes .
                                                                                                                                              Ikeno
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Pred. No. 9.6
0; Mismatches
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    prepared using DNA construct
d centromere containing copies

                                                                                                                                             Masumoto H,
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9.6e+03;
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                                                                                                                                             Okazaki T;
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This sequence

is a primer for a human alphoid DNA sequence that can

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The present invention describes the human Zmax1 gene and the high bone mass (HBM) gene, which are found on chromosome 11q13.3. The Zmax1 and HBM genes have osteopathic activities. The genes can be used in gene

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RESULT 1
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Best Local
                                                                                                                                                                                                                                                05-APR-2000;
05-APR-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human; high bone mass; HBM gene; 2max1 ge sequence tagged site; STS; osteoporosis; antisense therapy; vaccine; bone disorder
                                                              Disclosure;
                                                                                               New high bone mass (HBM) and modulating bone mass for the
                                                                                                                                                                                Carulli JP,
                                                                                                                                                                                                                                                                                                                                 18-OCT-2001.
                                                                                                                                                                                                                                                                                                                                                                  WO200177327-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                    sclerostosis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  HBM1 polymorphism containing amplicon PCR primer 1863F.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ABA82604;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ABA82604 standard; DNA; 18
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      generate transgenic mice.
                                                                                                                                                                                                                                                                                                 21-JUN-2000; 2000WO-US16951
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                                                                                                                                                                                                                 (GENO-) GENOME THERAPEUTICS CORP.
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1 ttctgagaa 9
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                                                                                                                                                                                                                                                                                                                                                                                                                   sapiens
                                                                                                                                                 2001-657171/75.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       BP; 3 A; 3 C; 4 G; 8 T; 0 other;
                                                                                                                                                                                Little
                                                                                                                                                                                                                                                  2000US-0543771.
2000US-0544398.
                                                                                                                                                                                                                                                                                                                                                                                                                                                    osteomalacia;
                                                              63; 443pp;
                                                                                                                                                                                  RD,
                                                                                                                                                                                  Recker RR,
                                                              English
                                                                                                                                                                                                                                                                                                                                                                                                                                                  fibrous dysplasia; PCR primer; linker; ss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  osteoporosis; osteopathic; gene therapy;
bone disorder; Paget's disease; adapter
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0;
                                                                                             treatment of e.g.
                                                                                                              Zmax1 genes and proteins useful for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 9;
Pred. No.
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                                                                                                                                                                                  Johnson
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     DB 19; 1
. 9.6e+03;
                                                                                                osteoporosis
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RESULT 13
ABA82646
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                       Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     sequence tagged site; STS; antisense therapy; vaccine; sclerostoe;
                                                                                                  The present invention describes the human Zmax1 gene and the high bone mass (HBM) gene, which are found on chromosome 11q.13.3. The Zmax1 and HBM genes have osteopathic activities. The genes can be used in gene therapy and in the production of vaccines. They can be used in the diagnosis and treatment of bone disorders including osteoporosis, Paget's disease, sclerostosis, osteomalacia and fibrous dysplasia. ABAB2708 to ABAB2700 and AAG68168 to AAG68193 represent sequences used in the exemplification of the present invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 therapy, antisense therapy and in the production of vaccines. They can be used in the diagnosis and treatment of bone disorders including osteoporosis, Paget's disease, solerostosis, osteomalacia and fibrous dysplasia. ABA82038 to ABA82700 and AAG68168 to AAG68193 represent
                                                                                                                                                                                                                                                                                                                                           05-APR-2000;
05-APR-2000;
                                                                                                                                                                                                            Disclosure;
                                                                                                                                                                                                                                                                      WPI;
                                                                                                                                                                                                                                                                                            Carulli JP,
                                                                                                                                                                                                                                                                                                                                                                             21-JUN-2000; 2000WO-US16951
                                                                                                                                                                                                                                                                                                                                                                                                                           WO200177327-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    sclerostosis; osteomalacia;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   25-JAN-2002
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                                                                               Sequence
                                                                                                                                                                                                                                  modulating
                                                                                                                                                                                                                                             New high bone mass (HBM)
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nes 9; Conserv
                                   Local
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                       Similarity
9; Conserv
                                                                                18
                                                                                                                                                                                                                                   bone mass
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PCR
                                                                               BP;
                                                                                                                                                                                                          Page 103; 443pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Conservative
                        Conservative
                                                                                                                                                                                                                                                                                                                                         2000US-0543771.
2000US-0544398.
                                                                                                                                                                                                                                                                                            Little
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            primer SEQ
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     the exemplification of the present invention.
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                                                                                                                                                                                                                                  (HBM) and for the
                                                                                                                                                                                                                                                                                           RD,
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                                                                                                                                                                                                                                                                                           Recker RR,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                gene; Zmaxl gene; chromosome 11; 11q13.3; osteoporosis; osteopath1c; gene therapy; bone disorder; Paget's disease; adapter; fibrous dysplasia; PCR primer; linker; ss.
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                     Score 9; DB
Pred. No. 9.6
0; Mismatches
                                                                                                                                                                                                                                  treatment of e.g. osteoporosis
                                                                                                                                                                                                                                             Zmax1 genes and proteins useful
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Pred. No.
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9.6e+03;
                                DB 22;
9.6e+03;
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                                           Length 18
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                       Indels
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Matches
                                                  Query Match
Best Local
                                                                                                                                                            A novel DNA construct comprises the present oligonucleotide (ON), which contains a core a regulatory element, operably linked to a promoter, which is operably linked to a heterologous gene (preferably a marker gene). The gene is under the transcriptional control of the promoter and the ON sequence when the ON is bound by a STAT protein activated in response to II-2, II-3, G-CSF, GM-CSF, erythropoietin, thrombopoietin, or preferably II-4, II-7, II-9, II-13 or II-15. Cytokine responsive host cells transfected with the DNA construct can be used to measure the ability of a compound to act as an agonist or antagonist of cytokine mediated gene transcription. In particular, they can be used to screen for cytokine mediated gene in the state of the state of protein and the state of protein cytokine modulators involved in the STATS and/or STAT6 protein a simplifier mathems.
                                                                                                                                                                                                                                                                                                                                                               DNA construct for screening modulators of cytokine-mediated transcription - contg. regulatory element and a cytokine-sensitive promoter operably linked to a heterologous gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                misc_feature
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              oligonucleotide;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              host cell; transfection; agonist; antagonist; mediated; transcription; modulation; signalling pathway; STAT6; oligonucleotide; electrophoretic mobility shift assay; E
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Oligonucleotide containing core DNA regulatory element.
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                                                                                                                                         signalling pathway. Following an electro
                                                                                                                                                                                                                                                                                                                                                                                                                     WPI; 1996-455362/45.
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                                                                                                  Sequence
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                                    Similarity
9; Conser
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                                                                                                   19
                                                                                                                           an electrophoretic to bind IL-4 and II
                                                                                                                                                                                                                                                                                                                                                                                                                                               Seidel
                                                                                                   BP;
                                      Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (first entry)
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                                                                                                                          ic mobility shift assay IL-13 moderately.
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Pred. No. 9.6
0; Mismatches
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                                                                                                                                         DNA construct
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RESULT 15
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ID AAV569
XX AAV569
XX Cytoki
KW Cytoki
KW transc
XX Synthe
PN US5814
XX 27-MAR
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PH LAMD I
XX WPI; 1
DNA CC
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PT TRANSC
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Search completed: August 20, 2002, 23:27:54 Job time: 3670 sec
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ANY56842-V56976 and ANV61601-V61631 are oligonucleotides used in the production of constructs comprising a cytokine-responsive regulatory celement linked to a promoter which is linked to a heterologous coding sequence so that the coding sequence is under the transcriptional control of the regulatory element and the promoter, where the regulatory celement has a nucleotide sequence selected from TTCNNGAA, TTANYTAA, and CTTCNYTAA where N is A, T, C or G, and y = 3 or 4. The constructs can be cused to detect or isolate transcription-regulating proteins, e.g. STAT proteins, in a sample by contacting the sample with the construct so that the protein binds to the regulatory element, and detecting or separating the resulting complex. The cells can be used in screening assays for coding sequence is measured in the presence and absence of a test very
                                                                                                                                                                                                                                                                                      Query Match
Best Local Similarity
Matches 9; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 19 BP; 4 A; 5 C; 4 G; 6 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Example 1; Column 19-20; 58pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              DNA constructs containing cytokine-responsive regulatory elements - useful in assays for transcription-regulating proteins or gene transcription agonists or antagonists
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Cytokine-responsive regulatory; primer; promoter; detection; isolation; transcriptional control; STAT protein; screening; agonist; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       02-DEC-1998 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAV56941 standard; DNA; 19 BP.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 27-MAR-1995;
14-APR-1994;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   27-MAR-1995;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAV56941;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (LIGA-) LIGAND PHARM INC.
                                                                                                                                                                      16 TTCTGAGAA 8
                                                                                                                                                                                                        1 ttctgagaa 9
                                                                                                                                                                                                                                                                                             100.0%;
ilarity 100.0%;
Conservative (
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94US-0228935.
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                                                                                                                                                                                                                                                                                                <u>,</u>
                                                                                                                                                                                                                                                                                             Score 9; DB 19;
Pred. No. 9.6e+03;
); Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                          Length 19;
                                                                                                                                                                                                                                                                                                Indels
                                                                                                                                                                                                                                                                                             0;
                                                                                                                                                                                                                                                                                         Gaps
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Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries
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Maximum DB seq length: 30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Total number of hits satisfying chosen parameters:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Scoring table:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Title:
Perfect score:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   OM nucleic -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score
             nucleic search, using sw model
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Issued_Patents_NA:*
1: /cgn2_6/ptodata/2/
2: /dgn2_6/ptodata/2/
3: /cgn2_6/ptodata/2/
4: /cgn2_6/ptodata/2/
5: /cgn2_6/ptodata/2/
6: /cgn2_6/ptodata/2/
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Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        IDENTITY_NUC Gapop 10.0 , Gapext 1.0
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Copyright (c) 1993 - 2000 Compugen Ltd
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       /cgn2_6/ptodata/2/ina/5A_COMB.seq:*
/cgn2_6/ptodata/2/ina/5B_COMB.seq:*
/cgn2_6/ptodata/2/ina/6A_COMB.seq:*
/cgn2_6/ptodata/2/ina/6B_COMB.seq:*
/cgn2_6/ptodata/2/ina/Backfiles1.seq:*
/cgn2_6/ptodata/2/ina/backfiles1.seq:*
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Length
           US-08-852-091-34
PCT-US95-04477-35
PCT-US95-04477-35
PCT-US95-04477-36
PCT-US95-684B-1819
US-08-585-684B-1820
US-09-038-073-1819
US-09-038-073-1820
US-09-411-020-56
US-08-411-07-9B-101
PCT-US95-04477-100
PCT-US95-04477-101
US-08-723-415B-21
US-09-078-311-7
US-08-600-982-20
US-09-189-627A-21
PCT-US94-11261A-20
US-08-564-955-18
US-08-564-955-18
US-08-564-955-18
US-08-564-955-18
US-08-564-955-18
US-08-564-955-18
US-08-564-955-18
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US-08-369-796-34
US-08-410-779B-35
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Sequence i3, Appl Sequence 34, Appl Sequence 34, Appl Sequence 34, Appl Sequence 34, Appl Sequence 1819, Appl Sequence 1820, App Sequence 1820, Appl Sequence 1820, Appl Sequence 100, Appl Sequence 100, Appl Sequence 101, Appl Sequence 101, Appl Sequence 101, Appl Sequence 21, Appl Sequence 18, Appl 
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Description
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US-08-411-020-13/c
; Sequence 13, Application US/08411020
; Patent No. 5712094
; Patent No. 5712094
                               Matches
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45	44	43	42	41	40	39	38	37	36	35	34	33	32	31	30	29	28
9	9	9	9	9	9	9	9	9	9	9	9	9	9	9	9	9	9
100.0	100.0	100.0	100.0	100.0	100.0	100.0	100.0	100.0	100.0	100.0	100.0	100.0	100.0	100.0	100.0	100.0	100.0
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US-08-379-078-663	US-08-379-078-558	US-08-379-078-553	US-09-026-039-70	US-08-912-272-70	US-08-781-891-141	US-08-781-891-39	US-09-159-417-4	US-08-953-041-4	US-09-067-284-2	US-09-590-778-18	US-09-501-698-18	US-09-165-060-18	US-09-133-508A-18	US-09-560-579A-13	US-09-232-863-18	US-09-099-015-18	US-09-075-511-18
Sequence 663, App	Sequence 558, App			70,	141,	39,	Sequence 4, Appli	Sequence 4, Appli	Sequence 2, Appli	Sequence 18, Appl	•	•	•	•	•	Sequence 18, Appl	`

ALIGNMENTS

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MOLECULE TYPE:
; DESCRIPTION:
; DESCRIPTION:
; US-08-411-020-13
                                                                                                                                                                                                                                                                                                                                              COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/411,020
FILING DATE: 27-MAR-1995
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: JUZGENSEN, Thomas E.
REGISTRATION NUMBER: 34,195
REGISTRATION NUMBER: 34,195
REGISTRATION NUMBER: 34,195
                                                                                                                                                                                 TELEFAX: (619) 535-3906
INFORMATION FOR SEQ ID NO: 13
SEQUENCE CHARACTERISTICS:
LENGTH: 9 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: SEIDEL, H. MARTI
APPLICANT: LAMB, I. PETER
APPLICANT: CHAN, SHIN-SHAY TIAN
TITLE OF INVENTION: METHODS AND ASSOCIATED REAGENTS FOR
TITLE OF INVENTION: DETECTING MODULATORS OF CYTOKINE ACTION
NUMBER OF SEQUENCES: 59
                                                                                                                                                                                                                                                                                                             TELECOMMUNICATION INFORMATION: TELEPHONE: (619) 550-7675
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      STREET: 9393 Tov
CITY: San Diego
STATE: Californ
                                                                                                                                                                    TOPOLOGY:
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    Similarity
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                                                                                                                                                                    linear
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                                                                                                   other nucleic acid
/desc = "OTHER NUCLEIC
SYNTHETIC DNA"
100.0%;
Score 9;
Pred. No.
DB 1; I
2.5e+07;
                 Length 9;
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                                                                                                                                                                        RESULT
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                                                                                                                    Sequence 35, Application US/08410779B Patent No. 5814517
                                                                                                                                                                                                                                                                                              Matches
                                                                                                                                                                                                                                                                                                            Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 34, Application US/08369796 Patent No. 5716622
                                                                                                    GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                             MOLECULE NO HYPOTHETICAL: NO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NAME: Jackson Esq., David A.
REGISTRATION NUMBER: 26,742
REFERENCE/DOCKET NUMBER: 600-1-116
TELECOMMUNICATION INFORMATION:
TELEPHONE: 201487-5800
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TITLE OF INVENTION: FÚNCTIONALLY ACTIVE REGIONS OF SIGNAL
TITLE OF INVENTION: TRANSDUCER AND ACTIVATOR OF TRANSCRIPTION (STAT) PROTEINS
NUMBER OF SEQUENCES:
9
CORRESPONDENCE ADDRESS:
ADDRESSEB: Klauber & Jackson
          APPLICANT: SEIDEL, H. MARTI
APPLICANT: LAMB, I. PETER
TITLE OF INVENTION: DNA SPACER REGULATORY ELEMENTS
TITLE OF INVENTION: RESPONSIVE TO CYTOKINES AND METHODS FOR THEIR USE
NUMBER OF SEQUENCES: 166
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: Jackson Esq., Davi
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: Zilong Wen
APPLICANT: Curt M. Horvath
APPLICANT: Zhong Zhong
CORRESPONDENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                   STRANDEDNESS:
TOPOLOGY: 11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TELEFAX:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICATION NUMBER: US/0 FILING DATE: 06-JAN-1995
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ENGTH:
                                                                                                                                                                                                                          TTCTGAGAA 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 nucleic acid
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                                                                                                                                                                                                                                                                                              Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                     linear
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ADDRESS
                                                                                                                                                                                                                                                                                                                                                                                                                                 DNA synthetic probe
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   double
                                                                                                                                                                                                                                                                                                            100.0%;
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Pred. No. 2.5
0; Mismatches
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2.5e+07
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ; DESCRIPTION: US-08-410-779B-35
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 34, Application US/08852091 Patent No. 5883228
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                GENERAL INFORMATION:
APPLICANT: James E. Darnell,
APPLICANT: Zilong Wen
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           REGISTRATION NUMBER: 34,195
REFERENCE/DOCKET NUMBER: 016-0013A.US
TELECOMMUNICATION INFORMATION:
TELEPHONE: (619) 550-7675
TELEFAX: (619) 535-3906
INFORMATION FOR SEQ ID NO: 35:
                                                          COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TITLE OF INVENTION: FUNCTIONALLY ACTIVE REGIONS OF SIGNAL TITLE OF INVENTION: TRANSDUCER AND ACTIVATOR OF TRANSCRIPTION (STAT) PROTEINS NUMBER OF SEQUENCES: 39
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOUTWARE: PATENTIN Release #1.0
CURRENT APPLICATION DATA:
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STREET: SAN DIEGO
CITY: CALIFORNIA
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LENGTH: 9 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: Curt M. Horapplicant: Zhong Zhong
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                                                                                                                                                                                                                                                                                                                                                                                                                  STREET: 411 Hack
CITY: Hackensack
                                                                                                                                                                                                                                                                                                      COUNTRY: UZIP: 07601
                                                                                                                                                                                                                                                                                                                                                                               STATE:
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TOPOLOGY: lir
APPLICATION NUMBER: US/08/852,091 FILING DATE: 06-MAY-1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ADDRESSEE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                     411 Hackensack Avenue
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/desc = "OTHER NUCLEIC ACID,
SYNTHETIC DNA"
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100.0%; Pred. No.
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PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/3
FILING DATE: 06-JAN-1995

08/369,796

CLASSIFICATION:

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                                           Query Match
Best Local Similarity
Thes 9; Conserve
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; Sequence 35, Application PC/TUS9504477
; GENERAL INFORMATION:
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                                                                                                                                        PCT-US95-04477-35
                                                                                                                                                                                                                                          PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08,
FILING DATE: 14-APR-1994
INFORMATION FOR SEQ ID NO: 35:
SEQUENCE CHARACTERISTICS:
LENGTH: 9 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
Best Local :
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TELEX: 133521
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
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                                                                                                                                                   MOLECULE TYPE: other nucleic acid
DESCRIPTION: /desc = "OTHER NUCLEIC ACID,
DESCRIPTION: SYNTHETIC DNA"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TITLE OF INVENTION: DNA SPACER REGULATORY ELEMENTS RESPONSIVE TO TITLE OF INVENTION: CYTOKINES AND METHODS FOR THEIR USE NUMBER OF SEQUENCES: 165
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                                                                                                                                                                                                                                                                                                                                                                                      MEDIUM TYPE: Floppy disk

MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC compatible

OPERATING SYSTEM: PC-DOS/MS-DOS

SOFTWARE: Patentin Release #1.0, Version #1.30 (EPO)
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                                                                                                                                                                                               STRANDEDNESS: sir
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                       APPLICATION NUMBER: PCT/US95/04477 FILING DATE:
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9 TTCTGAGAA 1
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nes 9; Conserva
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REFERENCE/DOCKET NUMBER: 600-1-116
                                                                                                                                                                                                                                                                                                                                          CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          STRANDEDNESS:
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9
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                                                                        100.0%; Score 9; DB 5; I
100.0%; Pred. No. 2.5e+07;
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Pred. No.
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                                                          Mismatches
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; ANTI-SENSE: NO
PCT-US95-17025-34
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PCT-US95-17025-34/c
                                                                                                                                                                                                          US-08-585-684B-1819/c
Patent NO. 30,,,,,
GENERAL INFORMATION:
GENERAL INFORMATION:
APPLICANT: Stinchcomb, Daniel T.
APPLICANT: Jarvis, Thale
APPLICANT: MCSWiggen, James
TITLE OF INVENTION: METHOD AND REAGENT FOR THE
TITLE OF INVENTION: INDUCTION OF GRAFT TOLERANCE
TITLE OF INVENTION: AND REVERSAL OF IMMUNE RESPONSES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 34, Application PC/TUS9517025 GENERAL INFORMATION:
                                                                                                                                                                          Sequence 1819, Application US/08585684B Patent No. 5877021
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                                                                                                                                                                                                                                                                                                                                                                                   Query Match
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INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
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TELEPHONE: 201 487-5800
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CURRENT APPLICATION DATA:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: Zilong Wen APPLICANT: Curt M. Horapplicant: Zhong Zhong
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TYPE: nucleic acid
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TELEFAX: 201 343-1684
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REGISTRATION NUMBER: 26,742
REFERENCE/DOCKET NUMBER: 600-1-116
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Curt M. Horvath
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Pred. No.
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CORRESPONDENCE ADDRESS:

ADDRESSEE: Lyon & Lyon STREET: 633 West Fifth Street STREET: Suite 4700

CITY: Los Angeles STATE: California

COUNTRY:

U.S.A.

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TELEX: 67-3510
INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS: LENGTH: 15 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
08-585-66.
                                                                                                                                                                                                                                                                                                                                                                                          Patent No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Matches
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Best Local S
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PRIOR APPLICATION DATA: 60/000,951
FILING DATE: July 7, 1995
ATTORNEY/AGENT INFORMATION:
NAME: Warburg, Richard
                                                           COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: storage
                                                                                                                                               CORRESPONDENCE ADDRESS:
ADDRESSEE: Lyon & Lyon
STREET: 633 West Fifth Street
STREET: Suite 4700
CITY: Los Angeles
STATE: California
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MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: storage
                                                                                                                                                                                                                                                                   TITLE OF INVENTION: METHOD AND REAGENT FOR THE TITLE OF INVENTION: INDUCTION OF GRAFT TOLERANCE AND REVERSAL OF IMMUNE RESPONSES
                                                                                                                                                                                                                                                                                                                 APPLICANT: Stinchcomb, Daniel T.
APPLICANT: Jarvis, Thale
APPLICANT: McSwigen, James
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TELECOMMUNICATION INFORMATION:
TELEPHONE: (213), 489-1600
TELEFAX: (213) 955-0440
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SOFTWARE: FastSEQ Version 1.5
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                         NUMBER OF SEQUENCES:
COMPUTER: IBM COMPATIBLE
OPERATING SYSTEM: IBM P.C. DOS 5.0
SOFTWARE: FastSEQ Version 1.5
JRRENT APPLICATION DATA:
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REGISTRATION NUMBER: 32,327
REFERENCE/DOCKET NUMBER: 21
                                                                                                                                COUNTRY:
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5. 5877021
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9; Conserv
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Conservative (
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Pred. No. 9.9
0; Mismatches
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 15 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
US-08-585-684B-1820
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Best Local Similarity
"---hes 9; Conserv?
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     GENERAL INFORMATION:
INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                    ZIP: 90071
COMPOTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: storage
COMPUTER: IBM COmpatible
OPERATING SYSTEM: IBM P.C. DOS 5.0
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TELECOMMUNICATION INFORMATION:
TELEPHONE: (213) 489-1600
TELEFAX: (213) 955-0440
TELEX: 67-3510
                                                                                                                                                 ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                 FILING DATE:
PRIOR APPLICATION DATA:
APPLICATION NUMBER:
                                                                                                                                                                                                                                                 SOFTWARE: FastSEQ Version 1.5
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICATION NUMBER: 60/00 FILING DATE: July 7, 1995 ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICATION NUMBER: US/08/585,684B FILING DATE: January 16, 1996 PRIOR APPLICATION DATA:
                                                                               TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NUMBER OF SEQUENCES:
                               TELEX:
                                              TELEFAX:
                                                                                                                NAME: Warburg, Richard REGISTRATION NUMBER: 3
                                                                                                                                                                     FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                   CITY: Los Angeles
STATE: California
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                                                               TELEPHONE:
                                                                                                  REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                    APPLICATION NUMBER:
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                                X: (213)
67-3510
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633 West Fifth Street
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Conservative 0;
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                                                (213) 489-1600
213) 955-0440
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                 1819:
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Pred. No.
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9.9e+02;
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; LENGTH: 15 base pairs
; TYPE: nucleic acid
; STRANDEDNESS: single
; TOPOLOGY: linear
US-09-038-073-1819
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US-09-038-073-1820
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                                                             Matches
                                                                                         Query Match
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                                                                                                                                                                                                                                  INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                               ATTORNEY/AGENT INFORMATION:
NAME: WAIDUTG, RICHARD
REGISTRATION UNMBER: 32,327
REFERENCE/DOCKET NUMBER: 218/
TELECOMMUNICATION INFORMATION:
TELEPHONE: (213) 489-1600
TELEFAX: (213) 955-0440
TELEX: 67-3510
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  GENERAL INFORMATION: APPLICANT: Stinch
                                                                                                                                                                                                  SEQUENCE CHARACTERISTICS:
LENGTH: 15 base pairs
                                                                                                                                                                                                                                                                                                                                                                        PRIOR APPLICATION DATA:
APPLICATION NUMBER:
FILING DATE:
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MEDIUM TYPE: 3.5" Diskette,
MEDIUM TYPE: storage
                                                                                                                                                                                                                                                                                                                                                                                                                                                    SOFTWARE: FastSEQ Version 1.5 CURRENT APPLICATION DATA:
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CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TITLE OF INVENTION: METHOD AND REAGENT FOR THE TITLE OF INVENTION: INDUCTION OF GRAFT TOLERANCE TITLE OF INVENTION: AND REVERSAL OF IMMUNE RESPONSES NUMBER OF SEQUENCES: 2751
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: Stinchcomb, Daniel T.
APPLICANT: Jarvis, Thale
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COMPUTER: IBM Compatible OPERATING SYSTEM: IBM P.C
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CITY: Los Angeles
STATE: California
                                                           Local Similarity
mes 9; Conserv
                                                                                                                                                                   TYPE: nucleic acid
STRANDEDNESS: sing
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TTCTGAGAA 4
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633 West Fifth Street
Suite 4700
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                                                             Conservative
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100.0%; Pred. No. 9.9
tive 0; Mismatches
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Pred. No.
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                                                           Mismatches
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9.9e+02;
hes 0; Indels
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RESULT 11
US-08-411-020-56/c
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                                                                                                                                                                                                                                    Sequence 57, Application US/08411020 Patent No. 5712094 GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Matches
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TELEGOMMUNICATION INFORMATION:
TELEPHONE: (619) 530-7675
TELEFAX: (619) 535-3906
INFORMATION FOR SEQ ID NO: 56:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICATION NUMBER: US/08/411
FILING DATE: 27-MAR-1995
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: JUTGENSEN, Thomas E.
REGISTRATION NUMBER: 34,195
REFERENCE/DOCKET NUMBER: 016-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    GENERAL INFORMATION:
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APPLICANT: LAMB, I. PETER
APPLICANT: CHAN, SHIN-SHAY TIAN
APPLICANT: CHAN, SHIN-SHAY TIAN
TITLE OF INVENTION: METHODS AND ASSOCIATED REAGENTS FOR
TITLE OF INVENTION: DETECTING MODULATORS OF CYTOKINE ACTION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         MOLECULE TYPE: other nucleic acid
DESCRIPTION: /desc = "OTHER NUCLEIC ACID,
DESCRIPTION: SYNTHETIC DNA"
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
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CORRESPONDENCE ADDRESSEE: Ligand Pharmaceuticals Incorporated STREET: 9393 Towne Centre Drive
                                                                                APPLICANT: SEIDEL, H. MARTI
APPLICANT: LAMB, I. PETER
APPLICANT: CHAN, SHIN-SHAY TIAN
APPLICANT: CHAN, SHIN-SHAY TIAN
APPLICANT: CHAN, SHIN-SHAY TIAN
APPLICANT: CHAN, SHIN-SHAY TIAN
TITLE OF INVENTION: DETECTING MODULATORS OF CYTOKINE ACT
NUMBER OF SEQUENCES: 59
CORRESPONDENCE ADDRESS:
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STRANDEDNESS: single
TOPOLOGY: linear
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CITY: San Diego
STATE: Californi
CITY: San Diego
STATE: California
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                                          ADDRESSEE: Ligand Pharmaceuticals Incorporated STREET: 9393 Towne Centre Drive
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Pred. No.
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1e+03;
                                                                                                                             OF CYTOKINE ACTION
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COMPUTER READABLE FORM: MEDIUM TYPE: Floppy

Floppy disk

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US-08-410-779B-100/c
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Best Local
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REGISTRATION NUMBER: 34,195
REFERENCE/DOCKET NUMBER: 016-0030.US
TELECOMMUNICATION INFORMATION:
TELEPHONE: (619) 550-7675
TELEPAX: (619) 535-3906
INFORMATION FOR SEQ ID NO: 57:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          GENERAL INFORMATION:
                                                                                                                                                                                                              COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
                                                PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/228,935
FILING DATE: 14-APR-1994
ATTORNEY/AGENT INFORMATION:
NAME: JURGENSEN, THOMAS E
REGISTRATION NUMBER: 34,195
FREGUSTRATION NUMBER: 34,195
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       MOLECULE TYPE: other nucleic acid
DESCRIPTION: /desc = "OTHER NUCLEIC ACID,
DESCRIPTION: SYNTHETIC DNA"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
         REGISTRATION NUMBER: 34,195
REFERENCE/DOCKET NUMBER: 016-0013A.US
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SEQUENCE CHARACTERISTICS:
LENGTH: 19 base pairs
TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: LAMB, I. PETER
TITLE OF INVENTION: DNA SPACER
NITILE OF INVENTION: RESPONSIVE
NUMBER OF SEQUENCES: 166
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: SEIDE APPLICANT: LAMB,
                                                                                                                                                                APPLICATION NUMBER: US/O FILING DATE: 27-MAR-1995 CLASSIFICATION: 435
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Local Similarity 100 hes 9; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                         3: LIGAND PHARMACEUTICALS INCORPORATED 9393 TOWNE CENTRE DRIVE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   linear
(619) 550-7675
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100.0%; Pr
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Pred. No. 1e+
0; Mismatches
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TO CYTOKINES AND METHODS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       DB 1; L., 1e+03; 0;
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US-08-410-7798-101
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Best Local Similarity
Thehes 9; Conserve
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; DESCRIPTION: /desc = "OTHER NUCLEIC ACID,
; DESCRIPTION: SYNTHETIC DNA"
US-08-410-779B-100
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 Query Match
Best Local S
Matches 9
                                                                                                                                                                                                                                                                                                                                                         COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PAtentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/410,779B
FILING DATE: 27-MAR-1995
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION UMBER: US 08/228,935
FILING DATE: 14-ARR-1994
ATTORNEY/AGENT INFORMATION:
NAME: JURGENSEN, THOMAS E
BEGITEMATION NUMBER: 34 105
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 101, Application US/08410779B Patent No. 5814517
                                                                                                                                                                                                                                                    REGISTRATION NUMBER: 34,195
REFERENCE/DOCKET NUMBER: 016-0013A.US
TELECOMMUNICATION INFORMATION:
TELEPHONE: (619) 550-7675
TELEFAX: (619) 535-3906
INFORMATION FOR SEQ ID NO: 101:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CORRESPONDENCE ADDRESS:
ADDRESSEE: LIGAND PHARMACEUTICALS INCORPORATED
STREET: 9393 TOWNE CENTRE DRIVE
                                                                                                                                              TOPOLOGY: 11
MOLECULE TYPE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SEQUENCE CHARACTERISTICS:
LENGTH: 19 base pairs
                                                                                                                                                                                                                                     SEQUENCE CHARACTERISTICS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CITY: SAN DIEGO
STATE: CALIFORNIA
                                                                                                                                                                            LENGTH: 19 base pairs TYPE: nucleic acid STRANDEDNESS: single
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  16 TTCTGAGAA 8
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h 100.0%; Score 9; DB 1; Similarity 100.0%; Pred. No. 1e+03; 9; Conservative 0; Mismatches
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LAMB, I. PETER
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/desc = "OTHER NUCLEIC ACID,
SYNTHETIC DNA"
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RESPONSIVE
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Title:
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Maximum Match 100%
Listing first 45 summaries
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Match
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em_htc:*
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AZ818898 ZMOO89N11
AZ313578 IMOO29N07
AZ303578 IMOO29N07
AZ303578 IMOO29N18
AZ313876 IMOO39N11
AZ387860 IMOO39N21
AZ387860 IMOO3517E15
AZ514561 IMO351N07
AZ5399Z20 IMO164P23
AZ810627 ZMOO76H17
AZ835621 ZMO179L21
AZ468141 IMO281504
AZ360212 IMO136V04
AZ36061 IMO215D17
AZ457101 IMO260317
AZ475949 T. brucei
AZ388663 IMO148J15
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qx13d05.x	2M0068B05	1M0407A03	1M0318M13	2M0102P15	2M0022B21	2M0018A16	1M0319H15	1M0321K14	1M0290A06	1M0354L24	2M0154P16	2M0089N11	T. brucei	1M0434B06	1M0082G16	AU012291	2M0014E09	1M0420K07	1M0361K01	MGS00966	T. brucei	1M0409E07	1M0367L12	1M0242F19	1M0098B15	1M0076A14	

ALIGNMENTS

	*																									
source	FEATURES							COMMENT	JOURNAL		TITLE			AUTHORS	REFERENCE		ORGANISM	SOURCE	KEYWORDS	VERSION	ACCESSION		DEFINITION	LOCUS	AZ818898	RESULT 1
1. 29 /organism="Mus musculus" /strain="C57BL/6J" /db_xref="taxon:10090" /clone="UUGC2M0089N11" /clone_1ib="Mouse 10kb plasmid UUGC1M library"	High quality sequence stop: 29. Location/Qualifiers	Plate: 0089 row: N column: 11 Seq primer: CGTTGTAAAACGACGGCCAGT Class: plasmid ends	Email: ddunn@genetics.utah.edu Insert Length: 10000 Std Error: 0.00	Fax: 801 585 7177	84112, USA Tel: 801 585 5606	Rm. 308, Biomedical Polymers Research Bldg., 20 S. 2030 E., SLC, UT	University of Utah	Contact: Robert B. Weiss		plasmid inserts	Mouse whole genome scaffolding with paired end reads from 10kb	,M., KOSE,M., KOSE,K., STOKES,K., Tingey,A., Von Niederhausern,A. and Wright.D.,Weiss.R.	Islam, H., Longacre, S., Mahmoud, M., Meenen, E., Pedersen, T., Reilly	Dunn, D., Aoyagi, A., Barber, M., Beacorn, T., Duval, B., Hamil, C.,	1 (bases 1 to 29)	Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus:	Mus musculus	house mouse.	GSS.	AZ818898.1 GI:12988806	AZ818898	clone UUGC2M0089N11 F, DNA sequence.	1F Mouse 10kb plasmid UUGC1M library Mus mus	AZ818898 29 bp DNA linear GSS 20-FFB-2001		

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REFERENCE
AUTHORS
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ORIGIN
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ORGANISM
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                                                                                                                                                                                                                                                                                                                                                                                                                                JOURNAL
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                                                                                                                                                            Email: ddunn@genetics.utah.edu
Insert Length: 10000 Sdd Error: 0.00
Plate: 0029 row: N column: 07
Seq primer: CACACAGGAAACAGCTATGACC
                                                                                                                                                                                                                                                                                                                                                                                                                              plasmid inserts
Unpublished (2000)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Dunn, D., Aoyagi, A., Barber, M., Beacorn, T., Duval, B., Hamil, C., Islam, H., Longacre, S., Mahmoud, M., Meenen, E., Pedersen, T., Reilly, M., Rose, M., Rose, R., Stokes, R., Tingey, A., von Niederhausern, A. and Wright, D., Weiss, R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus. 1 (bases 1 to 19)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                A2313531 19 bp DNA linear IM0029N07R Mouse 10kb plasmid UUGCIM library Mus clone UUGCIM0029N07 R, DNA sequence.
                                                                                         High quality sequence stop: 19.
Location/Qualifiers
                                                                                                                                                                                                                                                          Tel: 801 585 5606
Fax: 801 585 7177
                                                                                                                                                                                                                                                                                                                                                      University of Utah Genome Center University of Utah
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Mouse whole genome scaffolding with paired end reads from 10kb
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Mus musculus
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                                                                                                                                                                                                                                                                                                            USA
                                                                                                                                                                                                                                                                                                                               Biomedical Polymers Research Bldg.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 /sex="Male"
/lab_host="E. Coli strain XL10-Gold, T1-resistant, F-"
/note="Vector: PWD42nv; Purified genomic DNA from M.
musculus C57BL/6J (male) was obtained from the Jackson
Laboratory Mouse DNA Resource
/strain="C57BL/6J"
/db_xref="taxon:10090"
                                          organism-"Mus musculus"
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Pred. No.
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                                                                                                                                                                                                                                                                                            Contact: Robert B. W
University of Utah G
University of Utah
Rm. 308, Biomedical
                                                                                                         Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error: 0.00
Plate: 0003 row: H column: 07
Seq primer: CGTTGTNAAAACGACGGCCAGT
Class: plasmid ends
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Islam,H., Longacre,S., Mahmoud,M., Meenen,E., Pedersen,T., Reilly
,M., Rose,M., Rose,R., Stokes,R., Tingey,A., von Niederhausern,A.
and Wright,D.,Wells,R.
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                                                                                                                                                                                                                             Tel: 801 585 5606
Fax: 801 585 7177
                                                                                                                                                                                                                                                                                                                                                                                      Unpublished (2000)
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Mammalia; Eutheria; Rodentia; Sclurognathi; Muridae; Murinae; Mus
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polynucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pWN42 (gil4732114 [gb]AFI29072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chemically-competent E. coli XL10-Gold (Stratagene) cells and selected for ampicillin resistance."
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/organism="Mus musculus"
/strain="C57BL/6J"
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/sex="Male"
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100.0%; Pr
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%; Pred. No. 2.7
0; Mismatches
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o. 2.7e+05;
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                                                                                                                                                   Tel: 801 585 5606
Fax: 801 585 7177
Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error: (
Plate: 0091 row: H column: 18
Seg primer: CACACAGGAAACAGCTATGACC
                                                                                                                                                                                                                                                                                                                                                                                                                                                 plasmid inserts
Unpublished (2000)
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                                                                         High quality sequence stop: 24.
Location/Qualifiers
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and Wright,D.,Weiss,R.
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/organism="Mus musculus"
/strain="C57BL/6J"
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/note="Vector: PWD42nv; Purified genomic DNA from M.
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/clone_lib="Mouse 10kb plasmid UUGC1M library"
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                                                                                                                                           Tel: 801 585 5606
Fax: 801 585 7177
Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Errc
Plate: 0030 row: H column: 21
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Islam,H., Longacre,S., Mahmoud,M., Meenen,E., Pedersen,T., Reilly
,M., Rose,M., Rose,R., Stokes,R., Tingey,A., von Niederhausern,A.
and Wright,D.,Weiss,R.
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AZ313876.1 GI:10359208
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26 bp DNA linear GSS 29-SEP-200
1M0030H21F Mouse 10kb plasmid UUGC1M library Mus musculus genomic
clone UUGC1M0030H21 F, DNA sequence.
                                             High quality sequence stop: 26.
Location/Qualifiers
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/clone="UUGC2M0091H18"
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/organism≃"Mus musculus"
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/note="Vector: PWD42nv; Purified genomic DNA from M.
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100.0%; Pred. No. 2.8e+05;
^. Mismatches 0;
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                                      Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error: 0.00
Plate: 0147 row: M column: 20
Seq primer: CACACAGGAAACAGCTATGACC
Class: plasmid ends
High quality sequence. stop: 27.
                                                                                                                                                                                  Tel: 801 585 5606
Fax: 801 585 7177
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Unpublished (2000)
                                                                                                                                                                                                                                                                                                                                                                                                                  Dunn,D., Aoyagi,A., Barber,M., Beacorn,T., Duval,B., Hamil,C., Islam,H., Longacre,S., Mahmoud,M., Meenen,E., Pedersen,T., Reilly,M., Rose,M., Rose,R., Stokes,R., Tingey,A., von Niederhausern,A. and Wright,D.,Weiss,R.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1M0147M20R Mouse 10kb plasmid UUGC1M library Mus musculus genomic clone UUGC1M0147M20 R, DNA sequence.
A2387860
                                                                                                                                                                                                                                                                             University of Utah Genome Center University of Utah
                                                                                                                                                                                                                                                                                                                              Contact: Robert B.
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(http://www.jax.org/resources/documents/dnares/).
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/clone_lib="Mouse 10kb plasmid UUGC1M library"
/sex="Male"
                   Location/Qualifiers
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/db_xref="taxon:10090"
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pred. No. 2.8
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                  Tel: 801 585 5606

Fax: 801 585 7177

Email: ddunnegenetics.utah.edu
Insert Length: 10000 Std Error: 0.00
Plate: 0517 row: E column: 15
Seq primer: CGTTGTAAAACGACGGCCAGT
Class: plasmid ends
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Dunn,D., Aoyagi,A., Barber,M., Beacorn,T., Duval,B., Hamil,C.,

Islam,H., Longacre,S., Mahmoud,M., Meenen,E., Pedersen,T., Reilly,M., Rose,M., Rose,R., Stokes,R., Tingey,A., von Niederhausern,A. and Wright,D.,Weiss,R.
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Unpublished (2000)
High quality sequence stop: 28
                                                                                                                                                                                                                                                    University of Utah
                                                                                                                                                                                                                                                                                                           Contact: Robert B.
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/note="Vector: PWD42nv; Purified genomic DNA from M.
musculus C57BL/6J (male) was obtained from the Jackson
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/strain="C57BL/6J"
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/clone="UUGC1M0147M20"
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tive 0; Mismatches 0;
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Unpublished (2000)
Contact: Robert B. Weiss
University of Utah Genome C
University of Utah
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                                            Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error:
Plate: 0361 row: N column: 07
Seq primer: CGTTGTAAAACGACGGCCAGT
                                                                                                                                           Tel: 801 585 5606
Fax: 801 585 7177
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quality sequence stop: 30
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/db_xref="taxon:10090"
/clone="UUGC1M0517E15"
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/organism="Mus musculus"
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/note="Vector: PWD42nv; Purified genomic DNA from M.
musculus C57BL/6J (male) was obtained from the Jackson
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                       Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error:
Plate: 0.164 row: P column: 23
Seq primer: CACACAGGAAACAGCTATGACC
                                                                                                                                                                                                                                                                                                                     Mouse whole genome scaffolding with paired end reads from 10\mbox{kb} plasmid inserts
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                                                                                                                                                                                          University of Utah Genome Center
University of Utah
Rm. 308, Biomedical Polymers Research Bldg.,
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Contact: Robert B.
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Class: plasmid ends
                                                                                                                        Tel: 801 585 5606
Fax: 801 585 7177
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/clone_lib="Mouse 10kb plasmid UUGC1M library"
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                                                                                                                                                                    Contact: Robert B. Weiss
University of Utah Genome Center
University of Utah
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Dunn,D., Aoyagi,A., Barber,M., Beacorn,T., Duval,B., Hamil,C., Islam,H., Longacre,S., Mahmoud,M., Meenen,E., Pedersen,T., Reilly, M., Rose,M., Rose,R., Stokes,R., Tingey,A., von Niederhausern,A. and Wright,D.,Weiss,R.
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                                                                                  Tel: 801 585 5606
Fax: 801 585 7177
               Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error: 0.00
Plate: 0076 row: H column: 17
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musculus C57BL/6J (male) was obtained from t
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/clone="UUGC1M0164P23"
/clone_lib="Mouse 10kb plasmid UUGC1M library"
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/strain="C57BL/6J"
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High quality sequence stop:
Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Erro
Plate: 0129 row: L column: 2
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/strain="C57BL/6J"
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88.9%;
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                                                                                                            Contact: Robert B. Weiss
University of Utah Genome Center
University of Utah
Rm. 308, Biomedical Polymers Research Bldg., 20 S.
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                                             Tel: 801 585 5606
Fax: 801 585 7177
                                                                                                                                                                                                            plasmid inserts
Unpublished (2000)
                                                                                                                                                                                                                                                            Mouse whole genome scaffolding with paired end reads from 10kb
                                                                                                                                                                                                                                                                                     Dunn,D., Aoyagi,A., Barber,M., Beacorn,T., Duval,B., Hamil,C., Islam,H., Longacre,S., Mahmoud,M., Meenen,E., Pedersen,T., Reilly,M., Rose,M., Rose,R., Stokes,R., Tingey,A., von Niederhausern,A. and Wright,D.,Weiss,R.
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Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
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and selected for ampicillin resistance."
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/clone="UUGC2M0129L21"
/clone_lib="Mouse 10kb plasmid UUGC1M library"
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/strain="C57BL/6J"
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Pred. No. 7.2e+05;
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Seq primer: CGTTGTAAAACGACGGCCAGT
Class: plasmid ends
                                                                                 University of Utah University of Utah Rm. 308, Biomedical
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clone UUGC1M0103K04 F, DNA sequence.
                                                                                                                                                  Contact: Robert B.
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ddunn@genetics.utah.edu
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/note="Vector: PWD42nv; Purified genomic DNA from M.
musculus C57BL/6J (male) was obtained from the Jackson
Laboratory Mouse DNA Resource
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/clone="UUGC1M0281D04"
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/strain="C57BL/6J"
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Mammalla; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus

1 (bases 1 to 22)
                                                              Contact: Robert B. Weiss
University of Utah Genome Center
University of Utah
Rm. 308, Biomedical Polymers Research Bldg.,
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AZ430664
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/sex="Male"
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/strain="C57BL/6J"
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88.9%;
                                                                                                                                                                                                scaffolding with paired end reads from 10kb
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Pred. No. 7.3e+05;
0; Mismatches 1;
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Contact: Robert B. W
University of Utah G
University of Utah
Rui.
84112, ...
1: 801 F
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Bukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;

Bukaryota; Metazoa; Chordata; Sciurognathi; Muridae; Murinae; Mus

Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus

1 (bases 1 to 22)

Dunn,D., Aoyagi,A., Barber,M., Beacorn,T., Duval,B., Hamil,C.,

Islam,H., Longacre,S., Mahmoud,M., Meenen,E., Pedersen,T., Reilly

,M., Rose,M., Rose,R., Stokes,R., Tingey,A., von Niederhausern,A.

and Wright,D., Weiss,R.

Mouse whole genome scaffolding with paired end reads from 10kb
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            A2457101 22 bp DNA linear GSS 04-OCT-200 1M0260J17F Mouse 10kb plasmid UUGC1M library Mus musculus genomic clone UUGC1M0260J17 F, DNA sequence.
A2457101
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pWD42 (gil4732114|gb|AF129072.1), a copy number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chemically-competent E. coli xL10-Gold (Stratagene) cells and selected for ampicillin resistance."
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                                                             Biomedical
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/note="Vector: PWD42nv; Purified genomic DNA from M.
musculus C57BL/6J (male) was obtained from the Jack
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/strain="C57BL/6J"
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88.9%;
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Pred. No. 7.3e+05;
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Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error: 0.00
Plate: 0260 row: J column: 17
Seq primer: CGTTGTAAAACGACGGCCAGT
Class: plasmid ends
High quality sequence stop: 22.
Location/Qualifiers
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| Strains="Mus musculus" |
| Strain="C57BL/60" |
| /db_xref="taxon:10090" |
| Clone="UUGCIM0260J17" |
| /clone_lib="Mouse 10kb plasmid UUGCIM library" |
| /sex="Male" |
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